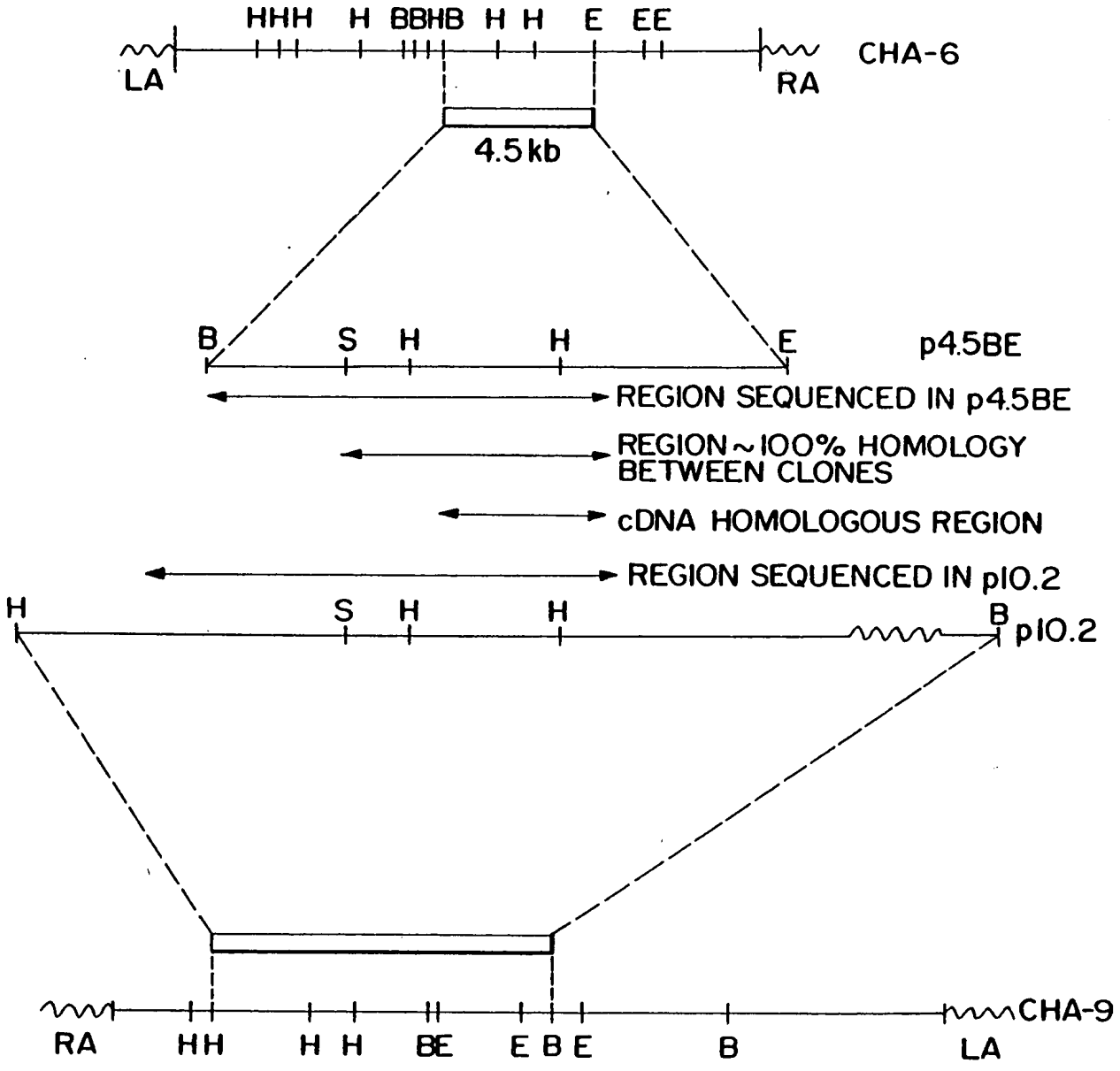
HHH H BBHB H H E EE
LA
RA
CHA-6
4.5kb
B S H H E
p4.5BE
REGION SEQUENCED IN p4.5BE
REGION ~100% HOMOLOGY BETWEEN CLONES
cDNA HOMOLOGOUS REGION
REGION SEQUENCED IN p10.2
H S H H B
p10.2
RA HH H H BE E B E B LA
CHA-9

FIG. 1